

Sequence Listing

<110> Gao, Wei-Qiang
Polakis, Paul
Shou, Jianyong
Smith, Victoria
Soriano, Robert
Williams, P. Mickey
Wu, Thomas D.
Zhang, Zemin

<120> COMPOSITIONS AND METHODS FOR THE DIAGNOSIS AND
TREATMENT OF TUMOR

<130> P5007R1-US

<150> 60/089,653

<151> 1998-06-17

<150> 60/090,355

<151> 1998-06-23

<150> 60/104,257

<151> 1998-10-14

<150> 60/119,537

<151> 1999-02-10

<150> 60/141,037

<151> 1999-06-23

<150> 60/145,698

<151> 1999-07-26

<150> 60/162,506

<151> 1999-10-29

<150> PCT/US99/12252

<151> 1999-06-02

<150> PCT/US99/20111

<151> 1999-09-01

<150> PCT/US99/28634

<151> 1999-12-01

<150> PCT/US99/28551

<151> 1999-12-02

<150> PCT/US00/00219

<151> 2000-01-05

<150> PCT/US00/00376

<151> 2000-01-06

<150> PCT/US00/04342

<151> 2000-02-18

<150> PCT/US00/08439
<151> 2000-03-30

<150> PCT/US00/13705
<151> 2000-05-17

<150> PCT/US00/23328
<151> 2000-08-24

<150> PCT/US01/06520
<151> 2001-02-28

<150> PCT/US01/20118
<151> 2001-06-22

<150> 09/888,257
<151> 2001-06-22

<160> 8

<210> 1
<211> 725
<212> DNA
<213> Homo Sapien

<400> 1
ccgcgggaac gctgtcctgg ctgcgcgcac ccgaacagcc tgtcctggtg 50
ccccggctcc ctgccccgcg ccagtcctg accctgcgcc cctcactcct 100
cccgtccat ctgctgctgc tgetgctget cagtgcggcg gtgtgccggg 150
ctgaggctgg gctcgaaacc gaaagtcccg tccggaccct ccaagtggag 200
accctggtgg agccccaga accatgtgcc gagcccgctg cttttggaga 250
cacgcttcac atacactaca cggaagctt ggtagatgga cgtattattg 300
acacctccct gaccagagac cctctgggta tagaacttgg ccaaagcag 350
gtgattccag gtctggagca gagtcttctc gacatgtgtg tgggagagaa 400
gcgaagggca atcattcctt ctcaactggc ctatggaaaa cggggatttc 450
caccatctgt ccagcggat gcagtgggtg agtatgacgt ggagctgatt 500
gcactaatcc gagccaaacta ctggctaaag ctggtgaagg gcattttgcc 550
tctggtaggg atggccatgg tgcagccct cctgggcctc attgggtatc 600
acctatacag aaaggccaat agacccaaag tctccaaaaa gaagctcaag 650
gaagagaaac gaaacaagag caaaaagaaa taataaataa taaattttaa 700
aaaacttaaa aaaaaaaaaa aaaaa 725

<210> 2
<211> 1174

<212> DNA
<213> Homo Sapien

<400> 2

gcggaactgg ctccggtgg cacctgagga gggcggtgac cccgagggcc 50
cagggagctg cccggtggc ctaggcaggc agccgcacca tggccagcac 100
ggccgtgcag cttctgggct tctgtctcag cttcctgggc atggtgggca 150
cgttgatcac caccatcctg ccgcactggc ggaggacagc gcacgtgggc 200
accaacatcc tcacggccgt gtcctacctg aaagggtctt ggatggagtg 250
tgtgtggcac agcacaggca tctaccagtg ccagatctac cgatccctgc 300
tggcgctgcc ccaagacctc caggctgccc gcgcctcat ggtcatctcc 350
tgcctgctct cgggcatagc ctgcgcctgc gccgtcatcg ggatgaagtg 400
cacgcgtgc gccaaaggga caccgcgcaa gaccaccttt gccatcctcg 450
gcggcaccct cttcatcctg gccggcctcc tgtgcatggt ggccgtctcc 500
tggaccacca acgacgtggt gcagaacttc tacaaccgcg tgcgtgccag 550
cggcatgaag tttgagattg gccaggccct gtacctgggc ttcattctct 600
cgtccctctc gctcattggt ggcacctgc tttgctgtc ctgccaggac 650
gaggcaccct acaggcccta ccaggccccg cccaggggcca ccacgaccac 700
tgcaaacacc gcacctgcct accagccacc agctgcctac aaagacaatc 750
gggccccctc agtgacctcg gccacgcaca gcgggtacag gctgaacgac 800
tacgtgtgag tccccacagc ctgcttctcc cctgggctgc tgtgggctgg 850
gtccccggcg ggactgtcaa tggaggcagg gggtccagca caaagtttac 900
ttctgggcaa tttttgtatc caaggaaata atgtgaatgc gaggaaatgt 950
ctttagagca cagggacaga gggggaaata agaggaggag aaagctctct 1000
ataccaaaga ctgaaaaaaaa aaatcctgtc tgtttttgta tttattatat 1050
atatttatgt gggtgatttg ataacaagtt taatataaag tgacttggga 1100
gtttggtcag tggggttggt ttgtgatcca ggaataaacc ttgcggatgt 1150
ggctgtttat gaaaaaaaaa aaaa 1174

<210> 3
<211> 3437
<212> DNA
<213> Homo Sapien

<400> 3

cagccccagc attgcagccg ctttcctgcc cacgttcatg tactgcctgg 2950
 gcagccagga ctttgaggtg gtgcagacgg ccctccggaa cctgcctgag 3000
 tacgtctctcc tgtgccaaga gcacgcggt gtgtctgtcc accgggcctt 3050
 cctggtgggc atgtacggcc agatggaccc cagcgcgcag atctccgagg 3100
 ccctgaggat cctgcatatg gaggcctga tgtgagcctg tggcagccga 3150
 cccccctcca agccccggcc cgtcccgtcc ccggggatcc tcgaggcaaa 3200
 gccaggaag cgtgggcgtt gctggtctgt ccgaggaggt gagggcgccg 3250
 agccctgagg ccaggcaggc ccaggagcaa tactccgagc cctgggggtg 3300
 ctccgggccg gccgctggca tcaggggccg tccagcaagc cctcattcac 3350
 cttctgggcc acagccctgc cgcggagcgg cggatcccc cgggcatggc 3400
 ctgggctggt tttgaatgaa acgacctgaa ctgtcaa 3437

<210> 4

<211> 1658

<212> DNA

<213> Homo Sapien

<400> 4

ggaaggcagc ggcagctcca ctcagccagt acccagatac gctgggaacc 50
 ttccccagcc atggcttccc tggggcagat cctcttctgg agcataatta 100
 gcatcatcat tattctggct ggagcaattg cactcatcat tggctttggt 150
 atttcagga gacactccat cacagtcact actgtcgct cagctgggaa 200
 cattggggag gatggaatcc tgagctgcac ttttgaacct gacatcaaac 250
 tttctgatat cgtgatacaa tggctgaagg aagggtgttt aggcttggtc 300
 catgagttca aagaaggcaa agatgagctg tcggagcagg atgaaatgtt 350
 cagaggccgg acagcagtgt ttgctgatca agtgatagtt ggcaatgcct 400
 ctttgcggt gaaaaacgtg caactcacag atgctggcac ctacaaatgt 450
 tatatcatca cttctaaagg caaggggaat gctaaccttg agtataaaac 500
 tggagccttc agcatgccg aagtgaatgt ggactataat gccagctcag 550
 agaccttgcg gtgtgaggct ccccgatggt tccccagcc cacagtggtc 600
 tgggcatccc aagttgacca gggagccaac ttctcggaag tctccaatac 650
 cagctttgag ctgaactctg agaatgtgac catgaagggt gtgtctgtgc 700
 tctacaatgt tacgatcaac aacacatact cctgtatgat tgaaaatgac 750

attgccaaag caacagggga tatcaaagt acagaatcgg agatcaaaag 800
 gcggagtcac ctacagctgc taaactcaaa ggcttctctg tgtgtctctt 850
 ctttctttgc catcagctgg gcacttctgc ctctcagccc ttacctgatg 900
 ctaaaataat gtgccttggc cacaaaaaag catgcaaagt cattgttaca 950
 acaggggatct acagaactat ttcaccacca gatatgacct agttttatat 1000
 ttctgggagg aaatgaattc atatctagaa gtctggagtg agcaaacaag 1050
 agcaagaaac aaaaagaagc caaaagcaga aggctccaat atgaacaaga 1100
 taaatctatc ttcaaagaca tattagaagt tgggaaaata attcatgtga 1150
 actagacaag tgtgttaaga gtgataagta aaatgcacgt ggagacaagt 1200
 gcatccccag atctcagggg cctccccctg cctgtcacct ggggagtgag 1250
 aggacaggat agtgcattgt ctttgtctct gaatttttag ttatatgtgc 1300
 tgtaatgttg ctctgaggaa gcccctggaa agtctatccc aacatatcca 1350
 catcttatat tccacaaatt aagctgtagt atgtacccta agacgctgct 1400
 aattgactgc cacttcgcaa ctccagggcg gctgcatttt agtaatgggt 1450
 caaatgattc actttttatg atgcttccaa aggtgccttg gcttctcttc 1500
 ccaactgaca aatgccaaag ttgagaaaaa tgatcataat tttagcataa 1550
 acagagcagt cggggacacc gattttataa ataaactgag caccttcttt 1600
 ttaaacaaaa aaaaaaaaaa aaaaaaaaaa aaaaaaaaaa aaaaaaaaaa 1650
 aaaaaaaaa 1658

<210> 5

<211> 201

<212> PRT

<213> Homo Sapien

<400> 5

Met	Thr	Leu	Arg	Pro	Ser	Leu	Leu	Pro	Leu	His	Leu	Leu	Leu	Leu
1				5				10					15	
Leu	Leu	Leu	Ser	Ala	Ala	Val	Cys	Arg	Ala	Glu	Ala	Gly	Leu	Glu
			20					25					30	
Thr	Glu	Ser	Pro	Val	Arg	Thr	Leu	Gln	Val	Glu	Thr	Leu	Val	Glu
			35					40					45	
Pro	Pro	Glu	Pro	Cys	Ala	Glu	Pro	Ala	Ala	Phe	Gly	Asp	Thr	Leu
			50					55					60	
His	Ile	His	Tyr	Thr	Gly	Ser	Leu	Val	Asp	Gly	Arg	Ile	Ile	Asp
			65					70					75	

Thr	Ser	Leu	Thr	Arg	Asp	Pro	Leu	Val	Ile	Glu	Leu	Gly	Gln	Lys	
				80					85					90	
Gln	Val	Ile	Pro	Gly	Leu	Glu	Gln	Ser	Leu	Leu	Asp	Met	Cys	Val	
				95					100					105	
Gly	Glu	Lys	Arg	Arg	Ala	Ile	Ile	Pro	Ser	His	Leu	Ala	Tyr	Gly	
				110					115					120	
Lys	Arg	Gly	Phe	Pro	Pro	Ser	Val	Pro	Ala	Asp	Ala	Val	Val	Gln	
				125					130					135	
Tyr	Asp	Val	Glu	Leu	Ile	Ala	Leu	Ile	Arg	Ala	Asn	Tyr	Trp	Leu	
				140					145					150	
Lys	Leu	Val	Lys	Gly	Ile	Leu	Pro	Leu	Val	Gly	Met	Ala	Met	Val	
				155					160					165	
Pro	Ala	Leu	Leu	Gly	Leu	Ile	Gly	Tyr	His	Leu	Tyr	Arg	Lys	Ala	
				170					175					180	
Asn	Arg	Pro	Lys	Val	Ser	Lys	Lys	Lys	Leu	Lys	Glu	Glu	Lys	Arg	
				185					190					195	
Asn	Lys	Ser	Lys	Lys	Lys										
				200											

<210> 6

<211> 239

<212> PRT

<213> Homo Sapien

<400> 6

Met	Ala	Ser	Thr	Ala	Val	Gln	Leu	Leu	Gly	Phe	Leu	Leu	Ser	Phe	
1				5					10					15	
Leu	Gly	Met	Val	Gly	Thr	Leu	Ile	Thr	Thr	Ile	Leu	Pro	His	Trp	
				20					25					30	
Arg	Arg	Thr	Ala	His	Val	Gly	Thr	Asn	Ile	Leu	Thr	Ala	Val	Ser	
				35					40					45	
Tyr	Leu	Lys	Gly	Leu	Trp	Met	Glu	Cys	Val	Trp	His	Ser	Thr	Gly	
				50					55					60	
Ile	Tyr	Gln	Cys	Gln	Ile	Tyr	Arg	Ser	Leu	Leu	Ala	Leu	Pro	Gln	
				65					70					75	
Asp	Leu	Gln	Ala	Ala	Arg	Ala	Leu	Met	Val	Ile	Ser	Cys	Leu	Leu	
				80					85					90	
Ser	Gly	Ile	Ala	Cys	Ala	Cys	Ala	Val	Ile	Gly	Met	Lys	Cys	Thr	
				95					100					105	
Arg	Cys	Ala	Lys	Gly	Thr	Pro	Ala	Lys	Thr	Thr	Phe	Ala	Ile	Leu	
				110					115					120	
Gly	Gly	Thr	Leu	Phe	Ile	Leu	Ala	Gly	Leu	Leu	Cys	Met	Val	Ala	

	125		130		135
Val Ser Trp Thr Thr Asn Asp Val Val Gln Asn Phe Tyr Asn Pro	140		145		150
Leu Leu Pro Ser Gly Met Lys Phe Glu Ile Gly Gln Ala Leu Tyr	155		160		165
Leu Gly Phe Ile Ser Ser Ser Leu Ser Leu Ile Gly Gly Thr Leu	170		175		180
Leu Cys Leu Ser Cys Gln Asp Glu Ala Pro Tyr Arg Pro Tyr Gln	185		190		195
Ala Pro Pro Arg Ala Thr Thr Thr Thr Ala Asn Thr Ala Pro Ala	200		205		210
Tyr Gln Pro Pro Ala Ala Tyr Lys Asp Asn Arg Ala Pro Ser Val	215		220		225
Thr Ser Ala Thr His Ser Gly Tyr Arg Leu Asn Asp Tyr Val	230		235		

<210> 7

<211> 1029

<212> PRT

<213> Homo Sapien

<400> 7

Met His Ile Leu Val Val His Ala Met Val Ile Leu Leu Thr Leu	1	5	10	15
Gly Pro Pro Arg Ala Asp Asp Ser Glu Phe Gln Ala Leu Leu Asp	20	25	30	
Ile Trp Phe Pro Glu Glu Lys Pro Leu Pro Thr Ala Phe Leu Val	35	40	45	
Asp Thr Ser Glu Glu Ala Leu Leu Leu Pro Asp Trp Leu Lys Leu	50	55	60	
Arg Met Ile Arg Ser Glu Val Leu Arg Leu Val Asp Ala Ala Leu	65	70	75	
Gln Asp Leu Glu Pro Gln Gln Leu Leu Leu Phe Val Gln Ser Phe	80	85	90	
Gly Ile Pro Val Ser Ser Met Ser Lys Leu Leu Gln Phe Leu Asp	95	100	105	
Gln Ala Val Ala His Asp Pro Gln Thr Leu Glu Gln Asn Ile Met	110	115	120	
Asp Lys Asn Tyr Met Ala His Leu Val Glu Val Gln His Glu Arg	125	130	135	
Gly Ala Ser Gly Gly Gln Thr Phe His Ser Leu Leu Thr Ala Ser	140	145	150	

Leu	Pro	Pro	Arg	Arg	Asp	Ser	Thr	Glu	Ala	Pro	Lys	Pro	Lys	Ser	155	160	165
Ser	Pro	Glu	Gln	Pro	Ile	Gly	Gln	Gly	Arg	Ile	Arg	Val	Gly	Thr	170	175	180
Gln	Leu	Arg	Val	Leu	Gly	Pro	Glu	Asp	Asp	Leu	Ala	Gly	Met	Phe	185	190	195
Leu	Gln	Ile	Phe	Pro	Leu	Ser	Pro	Asp	Pro	Arg	Trp	Gln	Ser	Ser	200	205	210
Ser	Pro	Arg	Pro	Val	Ala	Leu	Ala	Leu	Gln	Gln	Ala	Leu	Gly	Gln	215	220	225
Glu	Leu	Ala	Arg	Val	Val	Gln	Gly	Ser	Pro	Glu	Val	Pro	Gly	Ile	230	235	240
Thr	Val	Arg	Val	Leu	Gln	Ala	Leu	Ala	Thr	Leu	Leu	Ser	Ser	Pro	245	250	255
His	Gly	Gly	Ala	Leu	Val	Met	Ser	Met	His	Arg	Ser	His	Phe	Leu	260	265	270
Ala	Cys	Pro	Leu	Leu	Arg	Gln	Leu	Cys	Gln	Tyr	Gln	Arg	Cys	Val	275	280	285
Pro	Gln	Asp	Thr	Gly	Phe	Ser	Ser	Leu	Phe	Leu	Lys	Val	Leu	Leu	290	295	300
Gln	Met	Leu	Gln	Trp	Leu	Asp	Ser	Pro	Gly	Val	Glu	Gly	Gly	Pro	305	310	315
Leu	Arg	Ala	Gln	Leu	Arg	Met	Leu	Ala	Ser	Gln	Ala	Ser	Ala	Gly	320	325	330
Arg	Arg	Leu	Ser	Asp	Val	Arg	Gly	Gly	Leu	Leu	Arg	Leu	Ala	Glu	335	340	345
Ala	Leu	Ala	Phe	Arg	Gln	Asp	Leu	Glu	Val	Val	Ser	Ser	Thr	Val	350	355	360
Arg	Ala	Val	Ile	Ala	Thr	Leu	Arg	Ser	Gly	Glu	Gln	Cys	Ser	Val	365	370	375
Glu	Pro	Asp	Leu	Ile	Ser	Lys	Val	Leu	Gln	Gly	Leu	Ile	Glu	Val	380	385	390
Arg	Ser	Pro	His	Leu	Glu	Glu	Leu	Leu	Thr	Ala	Phe	Phe	Ser	Ala	395	400	405
Thr	Ala	Asp	Ala	Ala	Ser	Pro	Phe	Pro	Ala	Cys	Lys	Pro	Val	Val	410	415	420
Val	Val	Ser	Ser	Leu	Leu	Leu	Gln	Glu	Glu	Glu	Pro	Leu	Ala	Gly	425	430	435
Gly	Lys	Pro	Gly	Ala	Asp	Gly	Gly	Ser	Leu	Glu	Ala	Val	Arg	Leu			

	440	445	450
Gly Pro Ser Ser	Gly Leu Leu Val Asp	Trp Leu Glu Met Leu	Asp
	455	460	465
Pro Glu Val Val	Ser Ser Cys Pro Asp	Leu Gln Leu Arg Leu	Leu
	470	475	480
Phe Ser Arg Arg	Lys Gly Lys Gly Gln	Ala Gln Val Pro Ser	Phe
	485	490	495
Arg Pro Tyr Leu	Leu Thr Leu Phe Thr	His Gln Ser Ser Trp	Pro
	500	505	510
Thr Leu His Gln	Cys Ile Arg Val Leu	Leu Gly Lys Ser Arg	Glu
	515	520	525
Gln Arg Phe Asp	Pro Ser Ala Ser Leu	Asp Phe Leu Trp Ala	Cys
	530	535	540
Ile His Val Pro	Arg Ile Trp Gln Gly	Arg Asp Gln Arg Thr	Pro
	545	550	555
Gln Lys Arg Arg	Glu Glu Leu Val Leu	Arg Val Gln Gly Pro	Glu
	560	565	570
Leu Ile Ser Leu	Val Glu Leu Ile Leu	Ala Glu Ala Glu Thr	Arg
	575	580	585
Ser Gln Asp Gly	Asp Thr Ala Ala Cys	Ser Leu Ile Gln Ala	Arg
	590	595	600
Leu Pro Leu Leu	Leu Ser Cys Cys Cys	Gly Asp Asp Glu Ser	Val
	605	610	615
Arg Lys Val Thr	Glu His Leu Ser Gly	Cys Ile Gln Gln Trp	Gly
	620	625	630
Asp Ser Val Leu	Gly Arg Arg Cys Arg	Asp Leu Leu Leu Gln	Leu
	635	640	645
Tyr Leu Gln Arg	Pro Glu Leu Arg Val	Pro Val Pro Glu Val	Leu
	650	655	660
Leu His Ser Glu	Gly Ala Ala Ser Ser	Ser Val Cys Lys Leu	Asp
	665	670	675
Gly Leu Ile His	Arg Phe Ile Thr Leu	Leu Ala Asp Thr Ser	Asp
	680	685	690
Ser Arg Ala Leu	Glu Asn Arg Gly Ala	Asp Ala Ser Met Ala	Cys
	695	700	705
Arg Lys Leu Ala	Val Ala His Pro Leu	Leu Leu Leu Arg His	Leu
	710	715	720
Pro Met Ile Ala	Ala Leu Leu His Gly	Arg Thr His Leu Asn	Phe
	725	730	735

Gln	Glu	Phe	Arg	Gln	Gln	Asn	His	Leu	Ser	Cys	Phe	Leu	His	Val	740	745	750
Leu	Gly	Leu	Leu	Glu	Leu	Leu	Gln	Pro	His	Val	Phe	Arg	Ser	Glu	755	760	765
His	Gln	Gly	Ala	Leu	Trp	Asp	Cys	Leu	Leu	Ser	Phe	Ile	Arg	Leu	770	775	780
Leu	Leu	Asn	Tyr	Arg	Lys	Ser	Ser	Arg	His	Leu	Ala	Ala	Phe	Ile	785	790	795
Asn	Lys	Phe	Val	Gln	Phe	Ile	His	Lys	Tyr	Ile	Thr	Tyr	Asn	Ala	800	805	810
Pro	Ala	Ala	Ile	Ser	Phe	Leu	Gln	Lys	His	Ala	Asp	Pro	Leu	His	815	820	825
Asp	Leu	Ser	Phe	Asp	Asn	Ser	Asp	Leu	Val	Met	Leu	Lys	Ser	Leu	830	835	840
Leu	Ala	Gly	Leu	Ser	Leu	Pro	Ser	Arg	Asp	Asp	Arg	Thr	Asp	Arg	845	850	855
Gly	Leu	Asp	Glu	Glu	Gly	Glu	Glu	Glu	Ser	Ser	Ala	Gly	Ser	Leu	860	865	870
Pro	Leu	Val	Ser	Val	Ser	Leu	Phe	Thr	Pro	Leu	Thr	Ala	Ala	Glu	875	880	885
Met	Ala	Pro	Tyr	Met	Lys	Arg	Leu	Ser	Arg	Gly	Gln	Thr	Val	Glu	890	895	900
Asp	Leu	Leu	Glu	Val	Leu	Ser	Asp	Ile	Asp	Glu	Met	Ser	Arg	Arg	905	910	915
Arg	Pro	Glu	Ile	Leu	Ser	Phe	Phe	Ser	Thr	Asn	Leu	Gln	Arg	Leu	920	925	930
Met	Ser	Ser	Ala	Glu	Glu	Cys	Cys	Arg	Asn	Leu	Ala	Phe	Ser	Leu	935	940	945
Ala	Leu	Arg	Ser	Met	Gln	Asn	Ser	Pro	Ser	Ile	Ala	Ala	Ala	Phe	950	955	960
Leu	Pro	Thr	Phe	Met	Tyr	Cys	Leu	Gly	Ser	Gln	Asp	Phe	Glu	Val	965	970	975
Val	Gln	Thr	Ala	Leu	Arg	Asn	Leu	Pro	Glu	Tyr	Ala	Leu	Leu	Cys	980	985	990
Gln	Glu	His	Ala	Ala	Val	Leu	Leu	His	Arg	Ala	Phe	Leu	Val	Gly	995	1000	1005
Met	Tyr	Gly	Gln	Met	Asp	Pro	Ser	Ala	Gln	Ile	Ser	Glu	Ala	Leu	1010	1015	1020
Arg	Ile	Leu	His	Met	Glu	Ala	Val	Met									

<213> Homo Sapien

Thr Glu Ser Glu Ile Lys Arg Arg Ser His Leu Gln Leu Leu Asn
245 250 255

Ser	Lys	Ala	Ser	Leu	Cys	Val	Ser	Ser	Phe	Phe	Ala	Ile	Ser	Trp
				260					265					270
Ala	Leu	Leu	Pro	Leu	Ser	Pro	Tyr	Leu	Met	Leu	Lys			
				275					280					